

FIGURE 1A

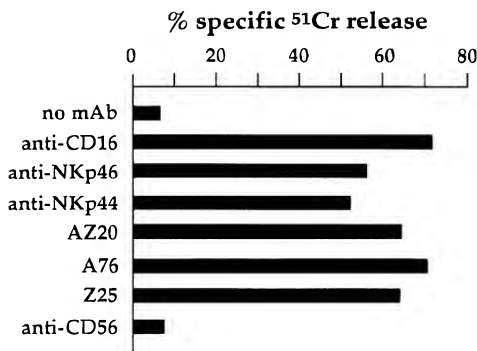


FIGURE 1B

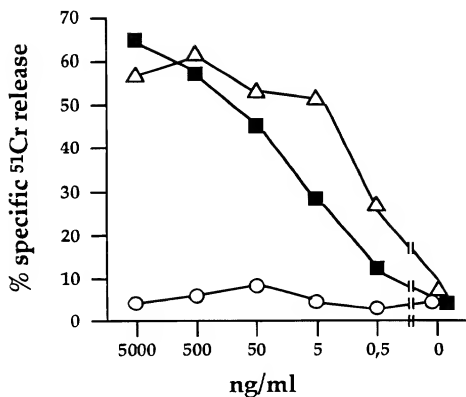


FIGURE 1C

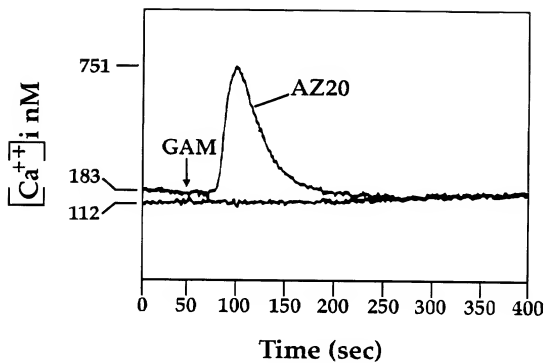


FIGURE 2A

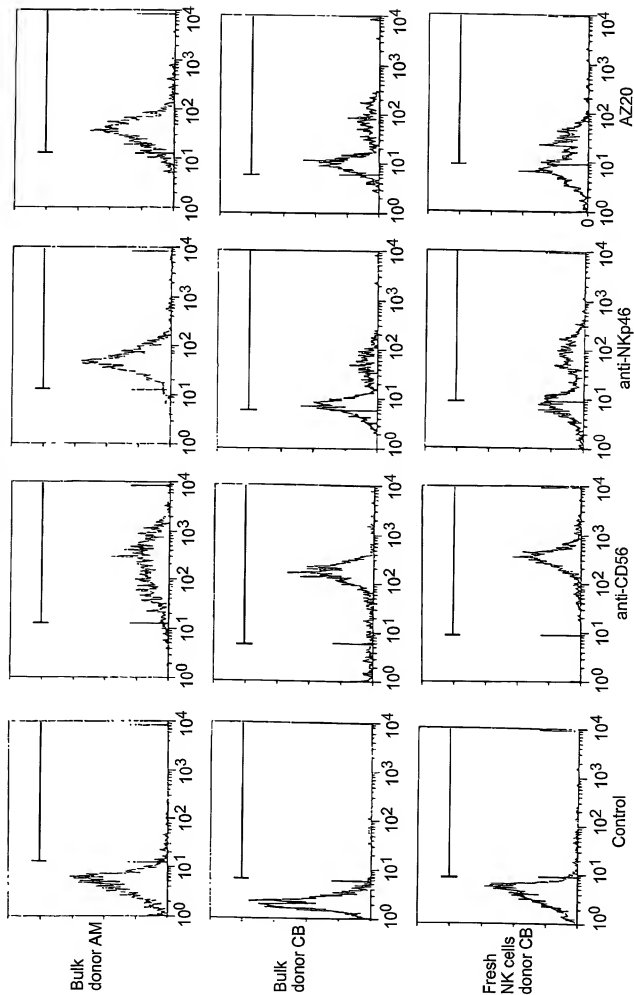


FIGURE 2B

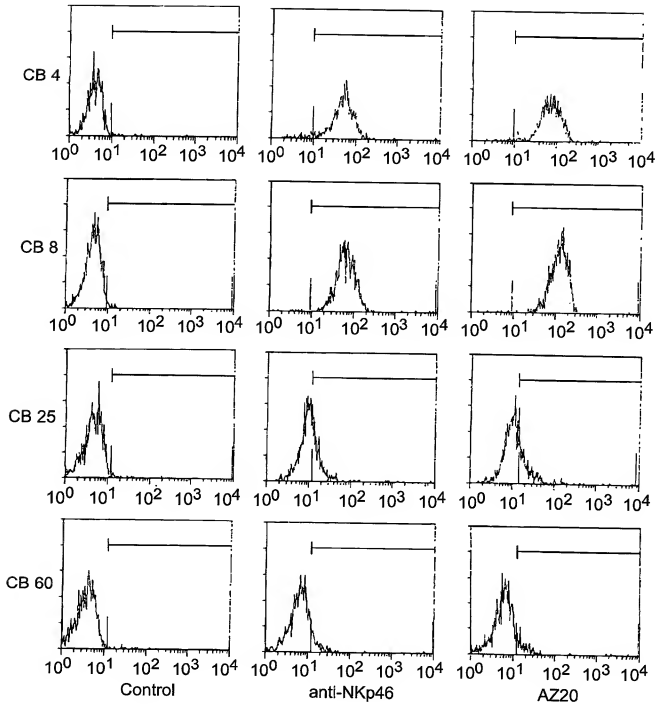


FIGURE 3A

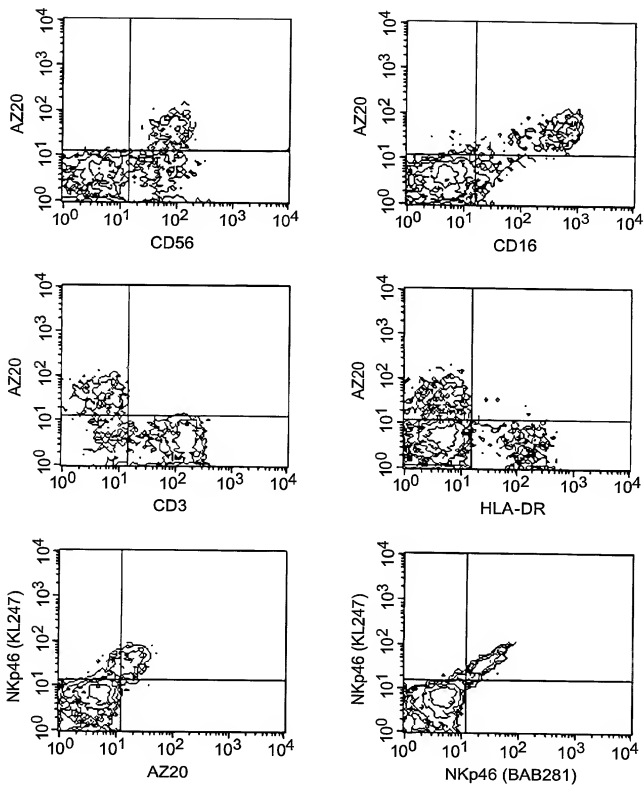


FIGURE 3B

Cells Daudi NK

69 —

45 —

28 —

18 —

Blot AZ20



FIGURE 4A

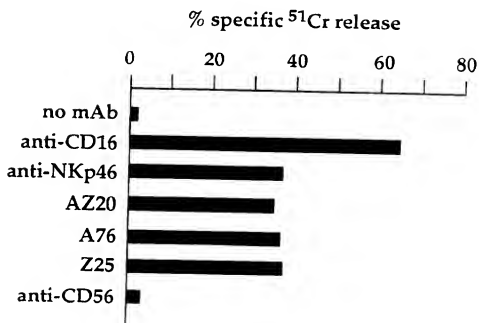


FIGURE 4B

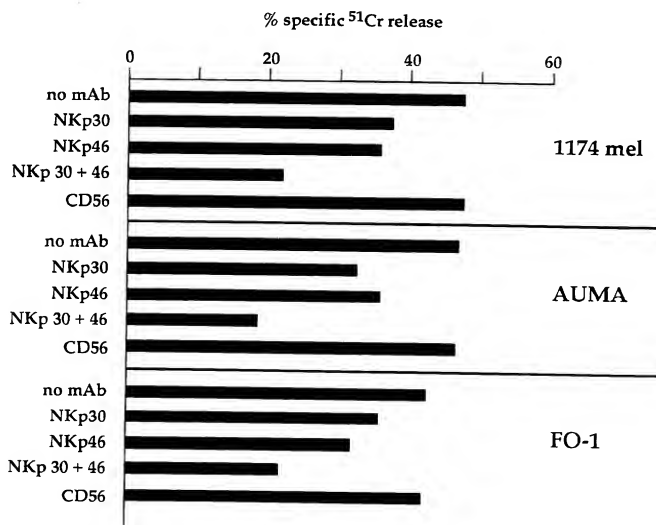


FIGURE 5

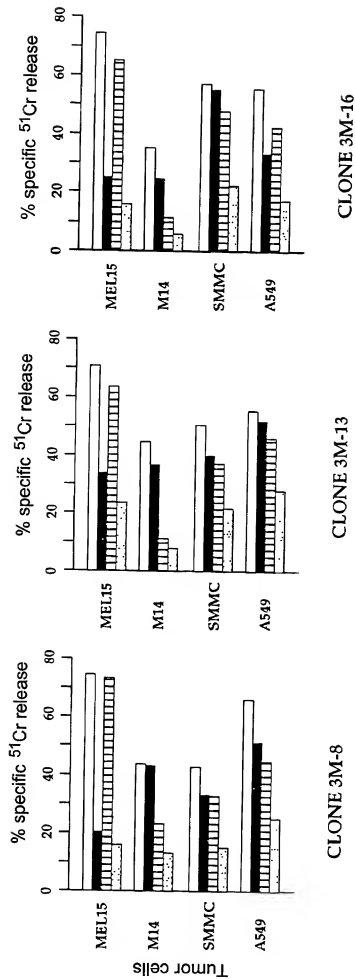


FIGURE 6A

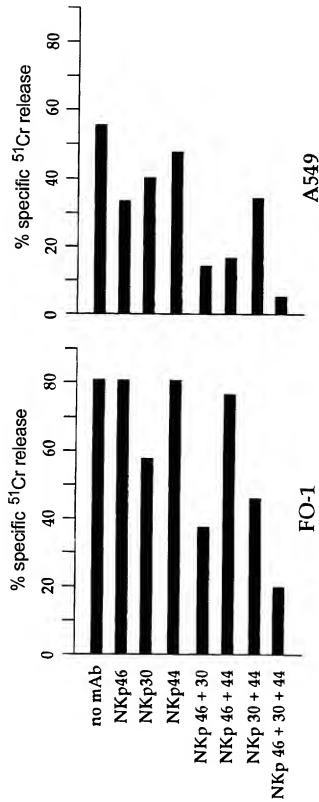


FIGURE 6B

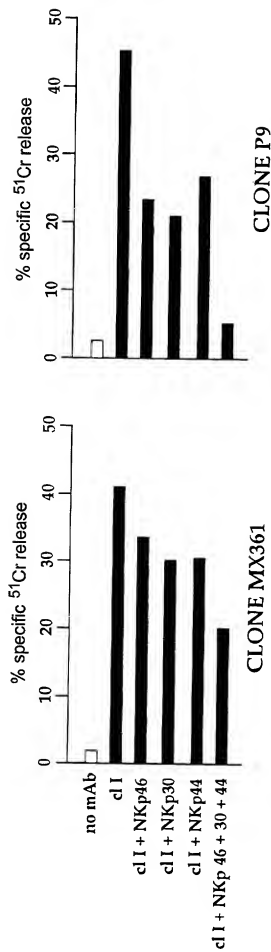


FIGURE 7A

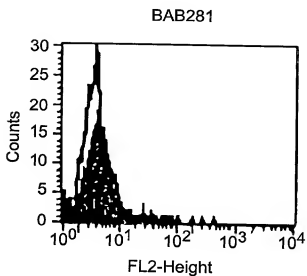
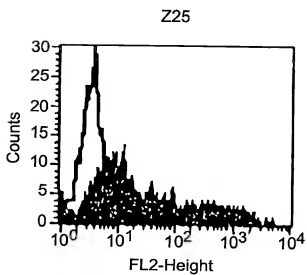
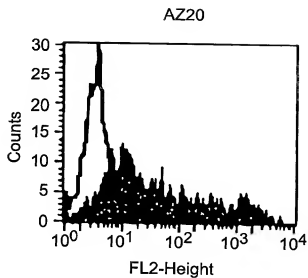
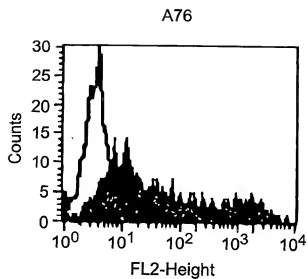


FIGURE 7B

SEQUENCE Nº 2

mcwmlllili	mvhpgscalw	VSQPEIRTL	EGSSAFLPQS	<u>FN</u> ASQGR LA I	50
GSVTWFRDEV	VPGKEVRNGT	PEFRGR LA PL	ASSRFLHDHQ	AELHIRDVRG	100
HDASIYVQRV	EVLGLGVGTG	<u>N</u> GTRLVVEKE	HPQLGAGTVL	<u>LLRAGFYAVS</u>	150
<u>FLSVAVGSTV</u>	YYQ GKCHCHM	GTHCHSSDGP	RGV IP EPRCP		190

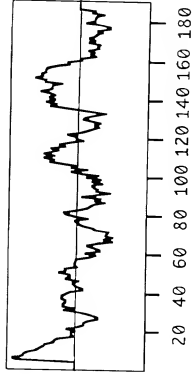


FIGURE 7C

SEQ ID NO 1

```

1  ccttcctctc  ccaccagac  ctcaactgtc  agatccctc  agccaaactg  gacatcttcc
61  gacatggcct  ggaatcctt  gctcaacttg  atcatgttcc  atccaggacc  cctgtctctc
121  tgggtgtccc  agcccccga  gattcgtacc  ctggaaggat  cctctgcttc  cctgccccgc
181  tccctcaatg  ccagccaagg  gagactggcc  attgccccg  tcaatgggtc  ccgagatgag
241  gtggttcacg  ggaaggaagt  gaggaatgga  acccaagatc  tcaggggccc  cctgccccca
301  ctctgtcttc  ccgtcttccc  ccataccacc  caggctgagc  tgcacatccc  ggaatgtcga
361  ggcctagacg  ccagcatcta  cgtgtgcaga  gtggagtctc  tggcccttgg  ctctgggaca
421  gggaaatggg  ctccggttgt  ggtggagaaa  gaacatccc  agctaggggc  tggtaacgtc
481  ctctctcttc  gggctggact  ctatgtcttc  agctttcttc  ctgtggccgt  gggcagcacc
541  gctctattac  agggcaaatg  ccactgtcac  atgggaacac  actgcccatt  cccagatcgg
601  ccccgaggcg  tgaattccaga  gccagatagt  cctctgtctc  ctccaaaaga  ccccaacaaa
661  tctgcccacc  cact

```

FIGURE 8A

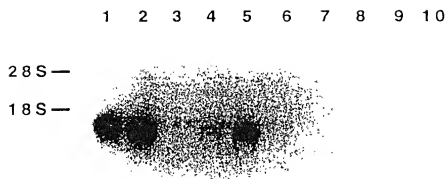


FIGURE 8B

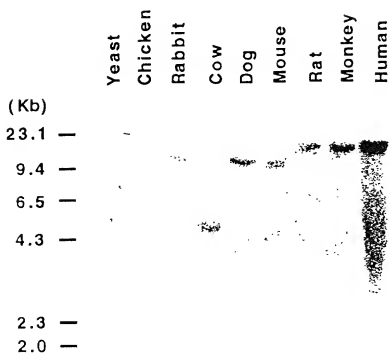


FIGURE 9A

Cells Daudi NK

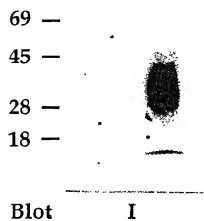


FIGURE 9B

